APPENDIX C-1 APPLICATION OF HARRINGTON ET AL. CLAIMS TO THE DISCLOSURE OF HARRINGTON ET AL. APPLICATION 08/941,223

| DISCLOSURE OF HARRINGTON ET AL. APPLICATION 08/941,223 | | |
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| | Harrington et al. Claim 271 | Harrington et al. Disclosure |
| | A method to activate expression of an endogenous gene in an isolated eukaryotic cell comprising | Abstract 7:23 8:9 24:20-21 30:3-10, 13-17, 27-28 31:8 32:19, 20, 22-25 Original claim 61 |
| | introducing a vector construct into said isolated eukaryotic cell, | Figures 1-4 Brief Description of the Figures 10:1-11:21 22:4-12 32:15-21 |
| | said vector construct comprising in operable combination | Figures 1-4 Brief Description of the Figures 10:1-11:21 9:24-25 17:21-18:2 19:1-21:6 25:17 26:9-23 |
| | 1) a promoter; | 10:14-15 |
| | 2) an exon sequence located 3' from and expressed by said promoter | Figures 1-4 Brief Description of the Figures 10:1-11:21 17:21-18:2 19:1-21:6 25:17 26:10-12 |
| | said exon being derived from a naturally occurring eukaryotic gene | Figure 1 25:17 -26:8 |
| | and not being a screenable marker gene; and | 25:30-26:2 26:30-27:2 28:14-16 |

28:24-27

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3) a splice donor sequence defining the 3' 26:2-3 region of said exon

said splice donor sequence being derived from a naturally-occurring eukaryotic gene; 27:4-9

wherein said vector construct is nonhomologously incorporated into the genome of a said isolated eukaryotic cell

12:5-21 14:29-15:24 15:28-16:4 27:12-14

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and said splice donor sequence of the transcript encoded by said exon is spliced to a splice acceptor sequence of said endogenous gene.

27:10-18